

Sequences producing significant alignments:

Score
(bits) E
Value

gi 37182174 gb AY358526.1 	Homo sapiens clone DNA66521 NXPH...	<u>3709</u>	0.0	
gi 21739859 emb AL834282.1 HSM805317	Homo sapiens mRNA; cDN...	<u>3701</u>	0.0	
gi 50481358 emb CR600551.1 	full-length cDNA clone CS0DF024...	<u>3693</u>	0.0	
gi 27754211 ref NM_007225.1 	Homo sapiens neurexophilin 3 (...)	<u>3681</u>	0.0	
gi 18490704 gb BC022541.1 	Homo sapiens neurexophilin 3, mR...	<u>3681</u>	0.0	
gi 28191449 gb AC006487.9 	Homo sapiens chromosome 17, clon...	<u>3489</u>	0.0	
gi 6330145 dbj AB032985.1 	Homo sapiens mRNA for KIAA1159 p...	<u>3420</u>	0.0	
gi 4105165 gb AF043468.1 AF043468	Homo sapiens neurexophilin 3 (...)	<u>1128</u>	0.0	
gi 4104960 gb AF042713.1 AF042713	Rattus norvegicus neurexophili...	<u>932</u>	0.0	
gi 11067402 ref NM_021679.1 	Rattus norvegicus neurexophilin 3 (...)	<u>932</u>	0.0	
gi 13938104 gb BC007167.1 	Mus musculus neurexophilin 3, mR...	<u>900</u>	0.0	
gi 18699993 ref NM_130858.1 	Mus musculus neurexophilin 3 (...)	<u>900</u>	0.0	
gi 26332926 dbj AK038955.1 	Mus musculus adult male hypotha...	<u>900</u>	0.0	
gi 21537471 emb AL662875.14 	Mouse DNA sequence from clone ...	<u>809</u>	0.0	
gi 17483894 gb AF424823.1 AF424823	Macaca mulatta neurexophilin 3 (...)	<u>533</u>	e-148	
gi 48841848 emb BX649458.8 	Zebrafish DNA sequence from clo...	<u>80</u>	2e-11	
gi 32251578 emb AL954329.7 	Zebrafish DNA sequence from clo...	<u>66</u>	3e-07	
gi 44844409 emb BX571820.5 	Zebrafish DNA sequence from clo...	<u>62</u>	5e-06	
gi 41635250 emb BX934722.1 	Gallus gallus finished cDNA, cl...	<u>62</u>	5e-06	
gi 49901442 gb BC076424.1 	Danio rerio cDNA clone MGC:10110...	<u>52</u>	0.004	
gi 49659361 emb BX323597.8 	Zebrafish DNA sequence from clo...	<u>50</u>	0.018	

gi 24980915 gb BC039747.1 	Mus musculus neurexophilin 3, mRNA	50	0.018	
gi 14198101 gb BC008114.1 	Mus musculus neurexophilin 3, mRNA	50	0.018	
gi 19068136 gb AY059393.1 	Mus musculus membrane glycoprotein	48	0.070	
gi 25046428 gb AC084073.22 	Mus musculus chromosome 1 clone	48	0.070	
gi 25013368 gb AC087781.17 	Mus musculus chromosome 1 clone	48	0.070	
gi 20988202 gb BC029659.1 	Mus musculus immunoglobulin superfamily	48	0.070	
gi 31543316 ref NM_053199.2 	Mus musculus immunoglobulin superfamily	48	0.070	
gi 26343202 dbj AK053077.1 	Mus musculus 15 days embryo head	48	0.070	
gi 26326786 dbj AK030782.1 	Mus musculus 8 days embryo whole body	48	0.070	
gi 40254885 ref NM_018990.2 	Homo sapiens chromosome X open reading frame	46	0.28	
gi 30354016 gb BC051881.1 	Homo sapiens chromosome X open reading frame	46	0.28	
gi 42600547 dbj AB162675.1 	Homo sapiens NXPH2 mRNA for neurexophilin 2	46	0.28	
gi 4678752 emb AL049683.1 HS753P092	Human gene from PAC 753	46	0.28	
gi 3550108 emb AL023653.1 HS753P9	Human DNA sequence from cDNA clone	46	0.28	
gi 15591370 emb AL589931.14 	Human DNA sequence from clone R	46	0.28	
gi 19718557 gb AF195953.2 	Homo sapiens membrane-bound amino acid transporter	46	0.28	
gi 15638787 gb AC092620.2 	Homo sapiens BAC clone RP11-231E11	46	0.28	
gi 4105163 gb AF043467.1 AF043467	Homo sapiens neurexophilin 2	46	0.28	
gi 42656448 ref XM_371573.2 	Homo sapiens neurexophilin 2 (mRNA)	46	0.28	
gi 11094302 dbj AB050725.1 	Bos taurus PREF1 gene, exon 1, mRNA	46	0.28	
gi 27545173 gb AC123069.4 	Mus musculus BAC clone RP23-23B2	44	1.1	
gi 3080664 gb AC004613.1 	Homo sapiens PAC clone RP4-651K2	44	1.1	
gi 48675466 gb AC144714.4 	Mus musculus BAC clone RP24-371M	44	1.1	
gi 28703719 gb BC047505.1 	Homo sapiens neurexophilin 1, mRNA	44	1.1	
gi 21754923 dbj AK095624.1 	Homo sapiens cDNA FLJ38305 first strand cDNA	44	1.1	
gi 12331138 emb AL445929.5 	Human DNA sequence from clone R	44	1.1	
gi 16303577 gb AC079239.6 	Homo sapiens BAC clone RP11-722M	44	1.1	
gi 18057083 gb AC093512.2 	Homo sapiens chromosome 16 clone	44	1.1	
gi 15741479 gb AC009014.5 	Homo sapiens chromosome 5 clone	44	1.1	
gi 6466800 gb AF113616.1 AF113616	Homo sapiens intestinal mRNA	44	1.1	
gi 4539679 gb AF073343.1 AF073343	Homo sapiens SMAD5 gene, mRNA	44	1.1	
gi 38229447 gb AC125536.3 	Mus musculus BAC clone RP24-142N	44	1.1	
gi 9929915 dbj AB038782.1 	Homo sapiens MUC3A gene for intestine	44	1.1	
gi 15823300 dbj AB047362.1 	Homo sapiens Nxph1 mRNA for neurexophilin 2	44	1.1	
gi 23097337 ref NM_152745.1 	Homo sapiens neurexophilin 1 (mRNA)	44	1.1	
gi 22759518 emb AL807794.6 	Mouse DNA sequence from clone R	44	1.1	
gi 39930314 ref NM_007224.1 	Homo sapiens neurexophilin 4 (mRNA)	42	4.3	
gi 37497210 gb AC126253.3 	Mus musculus BAC clone RP24-119M	42	4.3	
gi 31442528 gb AC127545.4 	Mus musculus BAC clone RP24-562H	42	4.3	
gi 34740425 gb AC122299.5 	Mus musculus BAC clone RP23-270L	42	4.3	
gi 30089839 gb AC133190.2 	Mus musculus BAC clone RP23-375H	42	4.3	
gi 38323217 gb AC110212.12 	Mus musculus chromosome 19, clone	42	4.3	
gi 29336308 gb AC127325.4 	Mus musculus BAC clone RP23-250E	42	4.3	
gi 21955088 gb AC122003.2 	Mus musculus BAC clone RP24-336G	42	4.3	
gi 37999311 gb AC118033.7 	Mus musculus chromosome 14, clone	42	4.3	
gi 37182349 gb AY358614.1 	Homo sapiens clone DNA83509 NXPH2	42	4.3	
gi 33469214 gb AC102604.13 	Mus musculus chromosome 19, clone	42	4.3	
gi 29791607 gb BC050314.1 	Homo sapiens bridging integrator	42	4.3	
gi 50548000 ref XM_501470.1 	Yarrowia lipolytica CLIB99, YAP1	42	4.3	
gi 31565572 gb BC053581.1 	Homo sapiens neurexophilin 4, mRNA	42	4.3	
gi 22003991 gb AC083905.20 	Homo sapiens 3 BAC RP11-124N24	42	4.3	

<u>gi 22830569 gb AF271897.2 </u>	Homo sapiens chromosome 8 clone ...	<u>42</u>	4.3	
<u>gi 47564193 gb AC131690.3 </u>	Mus musculus BAC clone RP23-386P...	<u>42</u>	4.3	
<u>gi 34530279 dbj AK124486.1 </u>	Homo sapiens cDNA FLJ42495 fis,...	<u>42</u>	4.3	
<u>gi 21753252 dbj AK094231.1 </u>	Homo sapiens cDNA FLJ36912 fis,...	<u>42</u>	4.3	
<u>gi 21750074 dbj AK091655.1 </u>	Homo sapiens cDNA FLJ34336 fis,...	<u>42</u>	4.3	
<u>gi 21321986 gb AC068657.6 </u>	Homo sapiens chromosome 5 clone ...	<u>42</u>	4.3	
<u>gi 46309689 gb AC147258.2 </u>	Mus musculus BAC clone RP24-297K...	<u>42</u>	4.3	
<u>gi 28875848 gb AC137834.9 </u>	Homo sapiens 12 BAC RP11-123K3 (...)	<u>42</u>	4.3	
<u>gi 20219181 gb AC103828.2 </u>	Homo sapiens chromosome 11, clon...	<u>42</u>	4.3	
<u>gi 27228898 gb AC037459.16 </u>	Homo sapiens chromosome 8, clon...	<u>42</u>	4.3	
<u>gi 14018353 emb AL355102.5 CNS05TCL </u>	Human chromosome 14 DNA...	<u>42</u>	4.3	
<u>gi 17483921 gb AF424833.1 AF424833 </u>	Macaca mulatta neurexoph...	<u>42</u>	4.3	
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<u>gi 32436442 emb AL773557.21 </u>	Mouse DNA sequence from clone ...	<u>42</u>	4.3	
<u>gi 13677161 gb AC012003.9 </u>	Homo sapiens BAC clone RP11-498P...	<u>42</u>	4.3	
<u>gi 14589684 gb AC021105.13 </u>	Homo sapiens BAC clone RP11-163...	<u>42</u>	4.3	
<u>gi 21627095 gb AE003806.3 </u>	Drosophila melanogaster chromoso...	<u>42</u>	4.3	
<u>gi 4105167 gb AF043469.1 AF043469 </u>	Homo sapiens neurexophilin ...	<u>42</u>	4.3	
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<u>gi 3047074 gb AF058914.1 F21E10 </u>	Arabidopsis thaliana BAC F2...	<u>42</u>	4.3	
<u>gi 11067404 ref NM_021680.1 </u>	Rattus norvegicus neurexophilin ...	<u>42</u>	4.3	
<u>gi 22798440 emb AL831725.5 </u>	Mouse DNA sequence from clone R...	<u>42</u>	4.3	
<u>gi 24395190 emb AL844583.6 </u>	Mouse DNA sequence from clone R...	<u>42</u>	4.3	
<u>gi 13092390 emb AL512294.3 LMFL654 </u>	Leishmania major Friedli...	<u>42</u>	4.3	
<u>gi 18252290 emb AL606504.14 </u>	Human DNA sequence from clone ...	<u>42</u>	4.3	
<u>gi 49647136 emb CR382129.1 </u>	Yarrowia lipolytica chromosome ...	<u>42</u>	4.3	
<u>gi 34304108 ref NM_183297.1 </u>	Mus musculus neurexophilin 4 (...)	<u>42</u>	4.3	
<u>gi 12835194 dbj AK004130.1 </u>	Mus musculus 18-day embryo whol...	<u>42</u>	4.3	

Alignments

Get selected sequences Select all Deselect all

>gi|37182174|gb|AY358526.1| Homo sapiens clone DNA66521 NXPH3 (UNQ687) mRNA, Length = 1871

Score = 3709 bits (1871), Expect = 0.0

Identities = 1871/1871 (100%)

Strand = Plus / Plus

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Query: 421 gtggaccatggcaatggacctt cagcgtccacttccaacacaatgccacaggccaaaggaa 480

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Figure 10.10. The effect of the number of hidden units on the performance of a neural network.

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For more information about the study, please contact Dr. John D. Cawley at (609) 258-4626 or via email at jdcawley@princeton.edu.

guttagen - 999-911-99999-99911-91951-1931

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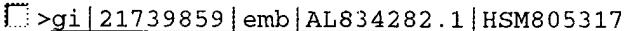
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 complete cds
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Score = 3701 bits (1867), Expect = 0.0
 Identities = 1870/1871 (99%)
 Strand = Plus / Plus

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Sbjct: 2109 aaccgctgattgctgactttgtgtgaagaatcgtgtttggagcagggaaataaagctt 2168

Query: 1861 gccccggggca 1871
Sbjct: 2169 gccccggggca 2179

[>] >gi|50481358|emb|CR600551.1| full-length cDNA clone CS0DF024YM05 of Fetal brain
(human)
Length = 2097

Score = 3693 bits (1863), Expect = 0.0
Identities = 1866/1867 (99%)
Strand = Plus / Plus

Query: 1 ctgggaccccggaaaagagaaggggagagcgagggacgagagcggaggaggaaatgcaa 60
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